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Correction



Correction: Complex Variation in Measures of General Intelligence and Cognitive Change

The PLOS ONE Staff

Figure 1 is missing panels B and C. The authors have provided a corrected version here. The publisher apologizes for the error.

Citation: The *PLOS ONE* Staff (2014) Correction: Complex Variation in Measures of General Intelligence and Cognitive Change. PLoS ONE 9(3): e91622. doi:10.1371/journal.pone.0091622

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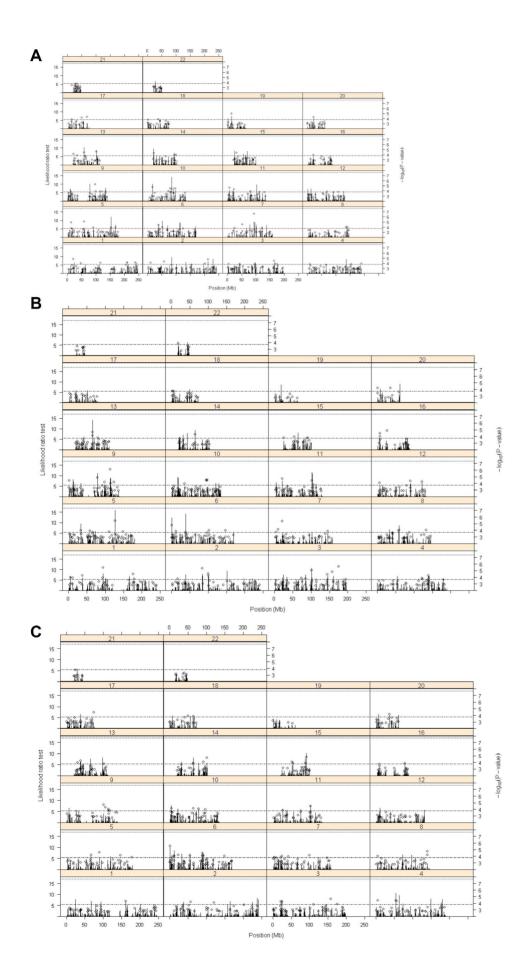


Figure 1. Plot of likelihood ratio test for phenotypic variance explained by each of 10,908 regions (groups of 101 consecutive SNPS) (bars) and $-\log_{10}$ P-values>2.7 for single SNP association (circles). Dashed line is 1% nominal significance threshold for LRT for individual regions, dotted line is 5% genome-wide significance threshold for individual regions obtained by permutation analysis. A crystallised intelligence n = 1791, B fluid intelligence n = 1706, and C cognitive change n = 1602. doi:10.1371/journal.pone.0081189.g001

Reference

 Rowe SJ, Rowlatt A, Davies G, Harris SE, Porteous DJ, et al. (2013) Complex Variation in Measures of General Intelligence and Cognitive Change. PLoS ONE 8(12): e81189. doi:10.1371/journal.pone.0081189